

5. The method according to claim 1, wherein said selected predetermined gDNA sequence within the 3'UTR or exon has a length of at least about 75 ~~to about 2000~~ bases/nucleotides.

Please add new claim 27, as follows:

27. The method according to claim 1, wherein said predetermined gDNA sequence within the 3'UTR or exon has a length of up to about 2000 nucleotides.